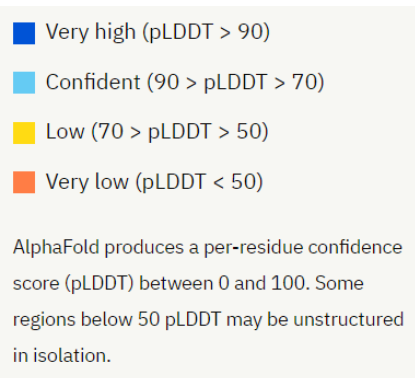


Supplementary material 2 – AlphaFold structures of predicted Z-DNA/Z-RNA binding proteins from *Homo sapiens*

Note: Model confidence legends depicted below, position of Z-alpha domain structural similarity is always highlighted in red dashed circle

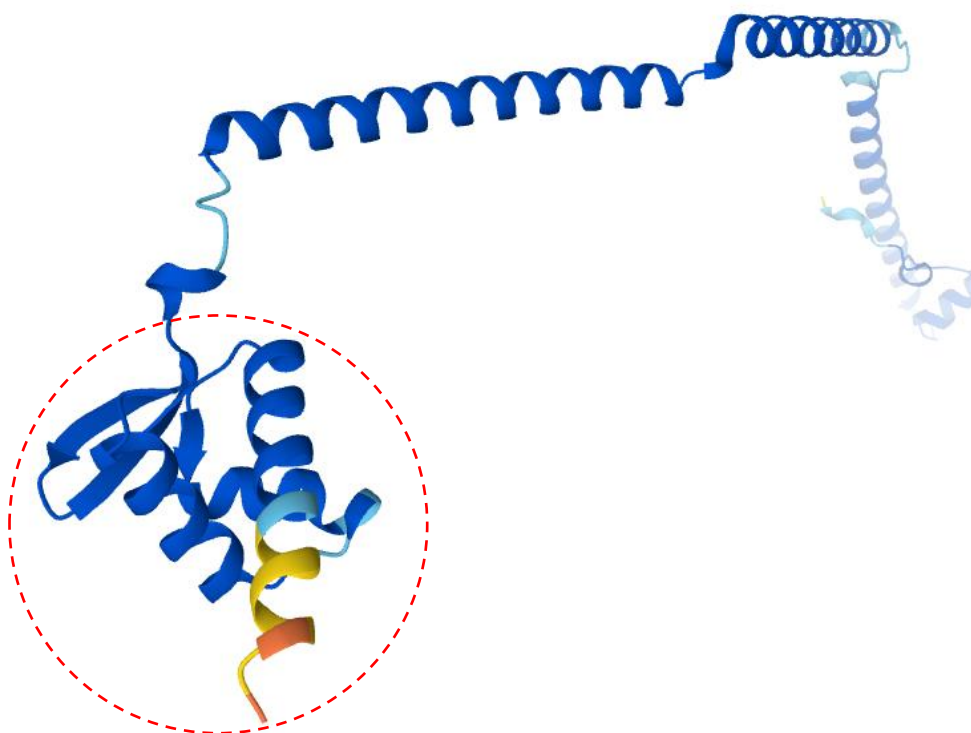


Accessed from: <https://alphafold.ebi.ac.uk/>

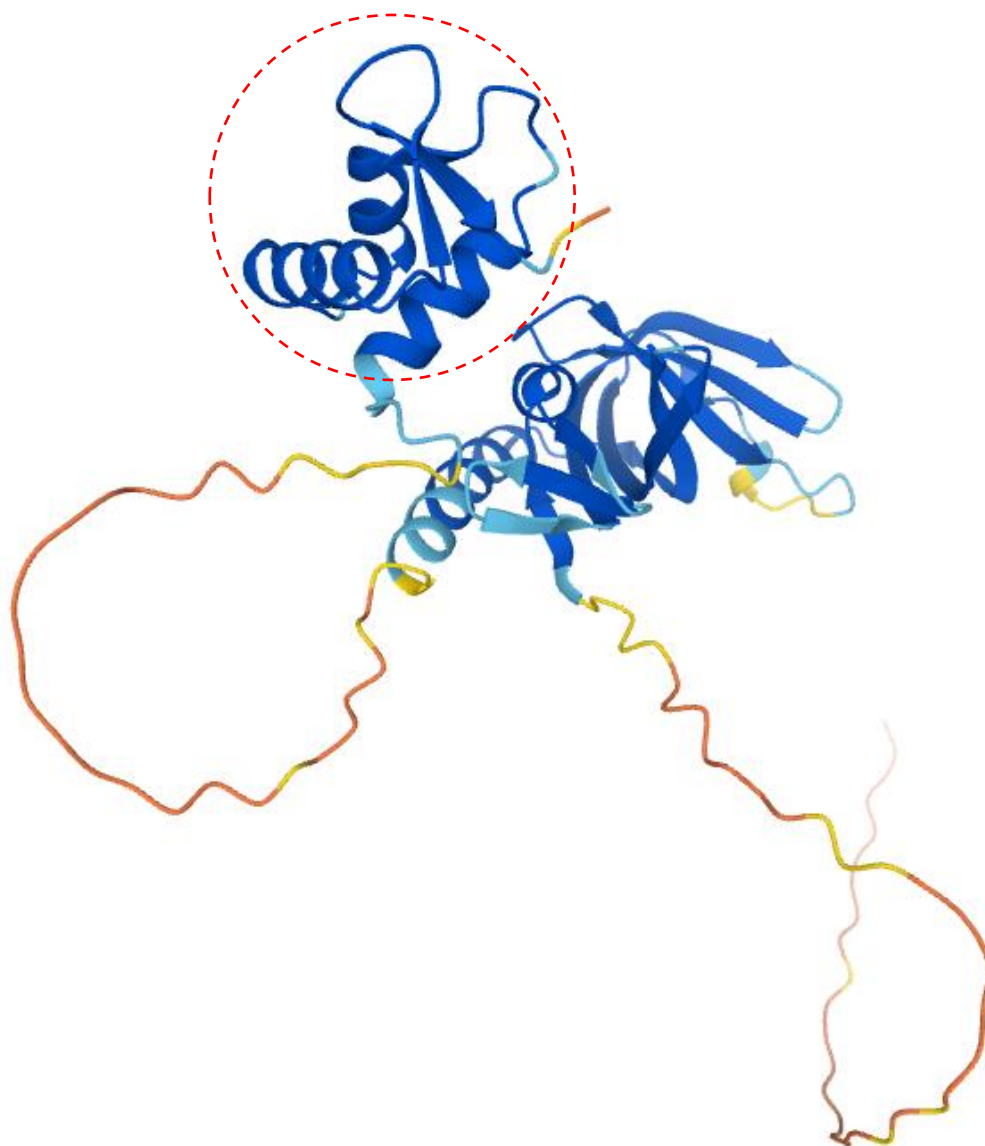
Citation:

Jumper, J et al. Highly accurate protein structure prediction with AlphaFold. Nature (2021).

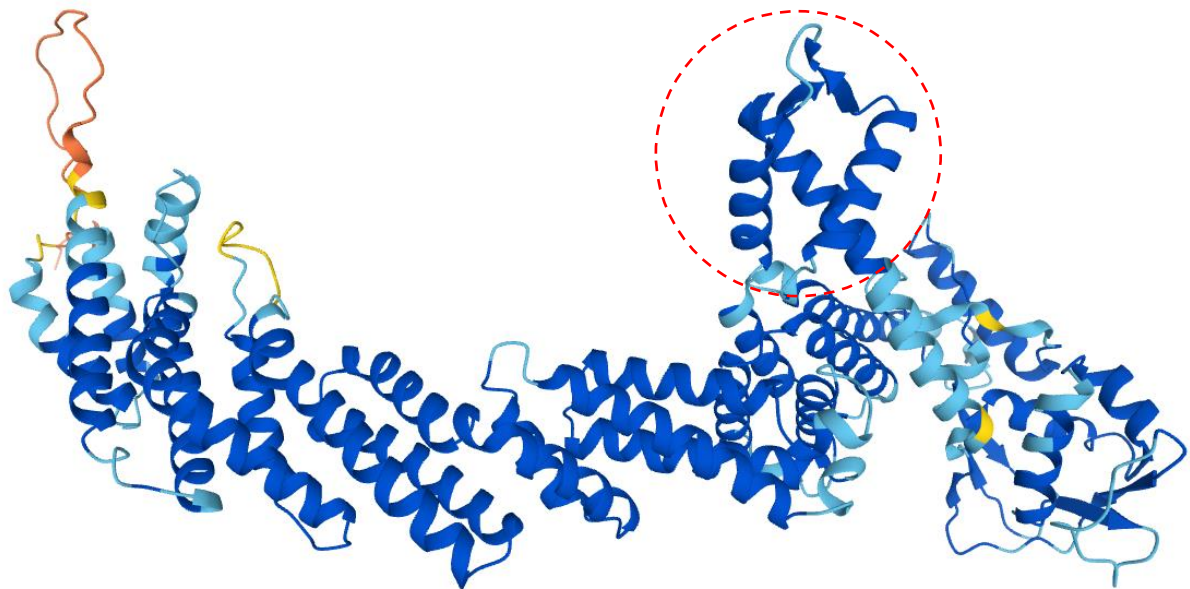
Homologous-pairing protein 2 homolog Q9P2W1 (HOP2_HUMAN)



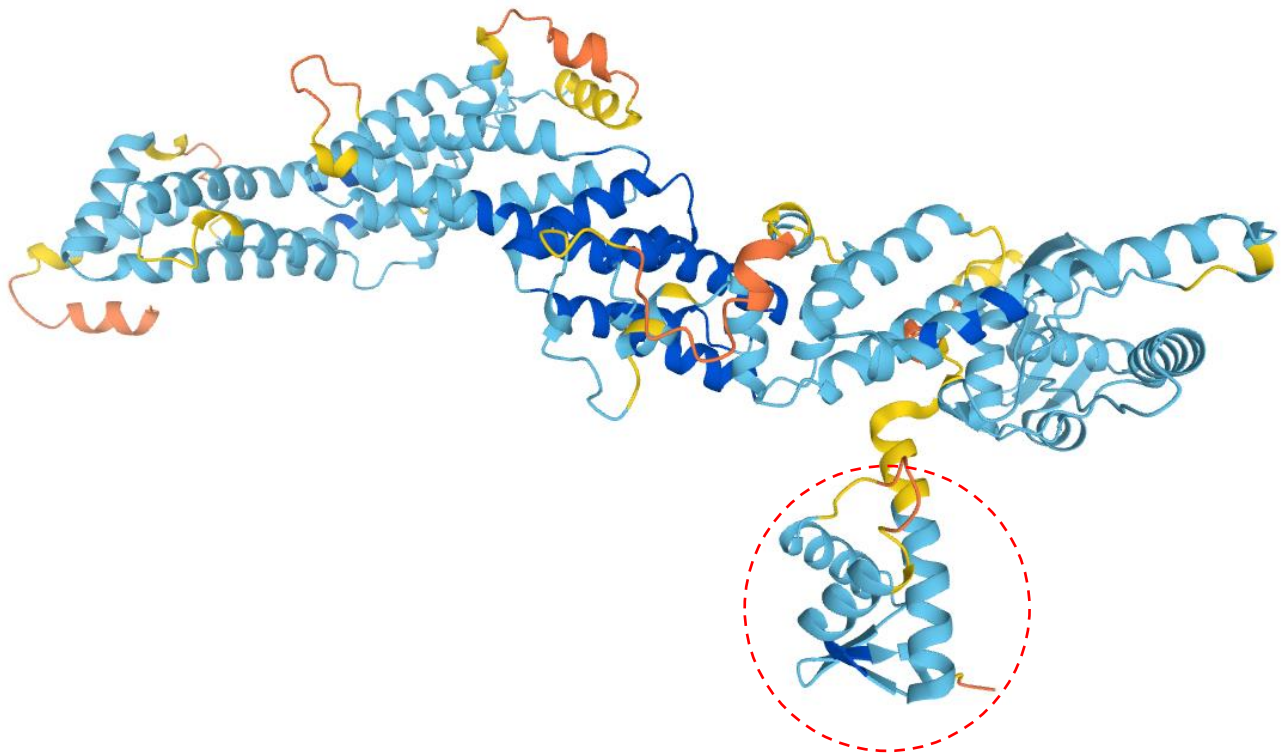
Replication protein A 32 kDa subunit P15927 (RFA2_HUMAN)



Cullin-1 Q13616 (CUL1_HUMAN)



Anaphase-promoting complex subunit 2 Q9UJX6 (ANC2_HUMAN)



DNA-directed RNA polymerase III subunit RPC6 Q9H1D9 (RPC6_HUMAN)

